

## Sequence Listing

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<120> METHOD FOR MAKING MONOCLONAL ANTIBODIES AND  
CROSS-REACTIVE ANTIBODIES OBTAINABLE BY THE METHOD

<130> P1468R1 (REVISED)

<140> US 09/329,633

<141> 1999-06-10

<150> US 60/089,253

<151> 1998-06-12

<160> 2

<210> 1

<211> 1799

<212> DNA

<213> human

<400> 1

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gcgccacaa aatacaccga cgatgcccga tctactttaa gggctgaaac 100  
ccacgggcct gagagactat aagagcgttc cctaccgcca tggaacaacg 150  
gggacagaac gccccggccg cttcgggggc ccggaaaagg cacggcccag 200  
gaccagggga ggcgcgggga gccaggcctg ggctccgggt cccaagacc 250  
cttgtgctcg ttgtcgccgc ggtcctgctg ttggtctcag ctgagtctgc 300  
tctgatcacc caacaagacc tagctcccca gcagagagcg gcccacaaac 350  
aaaagaggtc cagcccctca gagggattgt gtccacctgg acaccatata 400  
tcagaagacg gtagagattg catctcctgc aaatatggac aggactatag 450  
cactcactgg aatgacctcc ttttctgctt gcgctgcacc aggtgtgatt 500  
caggtgaagt ggagctaagt ccctgcacca cgaccagaaa cacagtgtgt 550  
cagtgcgaag aaggcacctt ccgggaagaa gattctcctg agatgtgccg 600  
gaagtgccgc acaggggtgc ccagagggat ggtcaaggtc ggtgattgta 650



<220>

<221> xaa

<222> 410

<223> xaa = leu or met

<400> 2

Met Glu Gln Arg Gly Gln Asn Ala Pro Ala Ala Ser Gly Ala Arg  
1 5 10 15

Lys Arg His Gly Pro Gly Pro Arg Glu Ala Arg Gly Ala Arg Pro  
20 25 30

Gly Leu Arg Val Pro Lys Thr Leu Val Leu Val Val Ala Ala Val  
35 40 45

Leu Leu Leu Val Ser Ala Glu Ser Ala Leu Ile Thr Gln Gln Asp  
50 55 60

Leu Ala Pro Gln Gln Arg Ala Ala Pro Gln Gln Lys Arg Ser Ser  
65 70 75

Pro Ser Glu Gly Leu Cys Pro Pro Gly His His Ile Ser Glu Asp  
80 85 90

Gly Arg Asp Cys Ile Ser Cys Lys Tyr Gly Gln Asp Tyr Ser Thr  
95 100 105

His Trp Asn Asp Leu Leu Phe Cys Leu Arg Cys Thr Arg Cys Asp  
110 115 120

Ser Gly Glu Val Glu Leu Ser Pro Cys Thr Thr Thr Arg Asn Thr  
125 130 135

Val Cys Gln Cys Glu Glu Gly Thr Phe Arg Glu Glu Asp Ser Pro  
140 145 150

Glu Met Cys Arg Lys Cys Arg Thr Gly Cys Pro Arg Gly Met Val  
155 160 165

Lys Val Gly Asp Cys Thr Pro Trp Ser Asp Ile Glu Cys Val His  
170 175 180

Lys Glu Ser Gly Ile Ile Ile Gly Val Thr Val Ala Ala Val Val  
185 190 195

Leu Ile Val Ala Val Phe Val Cys Lys Ser Leu Leu Trp Lys Lys  
200 205 210

Val Leu Pro Tyr Leu Lys Gly Ile Cys Ser Gly Gly Gly Gly Asp  
215 220 225

Pro Glu Arg Val Asp Arg Ser Ser Gln Arg Pro Gly Ala Glu Asp	230	235	240
Asn Val Leu Asn Glu Ile Val Ser Ile Leu Gln Pro Thr Gln Val	245	250	255
Pro Glu Gln Glu Met Glu Val Gln Glu Pro Ala Glu Pro Thr Gly	260	265	270
Val Asn Met Leu Ser Pro Gly Glu Ser Glu His Leu Leu Glu Pro	275	280	285
Ala Glu Ala Glu Arg Ser Gln Arg Arg Arg Leu Leu Val Pro Ala	290	295	300
Asn Glu Gly Asp Pro Thr Glu Thr Leu Arg Gln Cys Phe Asp Asp	305	310	315
Phe Ala Asp Leu Val Pro Phe Asp Ser Trp Glu Pro Leu Met Arg	320	325	330
Lys Leu Gly Leu Met Asp Asn Glu Ile Lys Val Ala Lys Ala Glu	335	340	345
Ala Ala Gly His Arg Asp Thr Leu Tyr Thr Met Leu Ile Lys Trp	350	355	360
Val Asn Lys Thr Gly Arg Asp Ala Ser Val His Thr Leu Leu Asp	365	370	375
Ala Leu Glu Thr Leu Gly Glu Arg Leu Ala Lys Gln Lys Ile Glu	380	385	390
Asp His Leu Leu Ser Ser Gly Lys Phe Met Tyr Leu Glu Gly Asn	395	400	405
Ala Asp Ser Ala Xaa Ser	410		